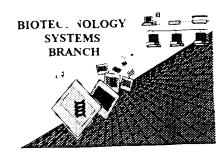
RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/94/,/99
Source: 0/ff

Source:

Date Processed by STIC:

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS. PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST 25

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2Kcompliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/941,179

DATE: 09/18/2001 TIME: 11:34:59

pg 1-2

Input Set : A:\sequence-text.txt

Output Set: N:\CRF3\09182001\I941179.raw

3 <110> APPLICANT: Bayer Aktiengesellschaft

5 <120> TITLE OF INVENTION: Acetylcholine receptor subunits

7 < 130 > FILE REFERENCE: Le A 34 821

9 <140 > CURRENT APPLICATION NUMBER: US/09/941,179 10 <141 > CURRENT FILING DATE: 2001-08-27

12 <150> PRIOR APPLICATION NUMBER: DE 100 42 177.6

13 <151> PRIOR FILING DATE: 2000-08-28

15 <160> NUMBER OF SEQ ID NOS: 17

17 <170> SOFTWARE: PatentIn Ver. 2.1

Does Not Comply Corrected Diskette Needed

ERRORED SEQUENCES

1056 <210> SEQ ID NO: 17

1057 <211> LENGTH: 75

1058 <212> TYPE: DNA

1059 <213> ORGANISM: Artificial Sequence

1061 <220> FEATURE:

1062 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer

1064 <400> SEQUENCE: 17

1065 ggtagggctc gtcgcagcat gtataaaact tttcgttgcg aacggccggg acttcaatga 60

E--> 1066 tgacccactc cccgc

E - - > 1068 (1)

4/941,179 <u>2</u>

<210><211><212><212><213><223>	622 Line 1 / 1707 Likeway	12227, 62227, 62227
<400>	receptor	The show

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/941,179

DATE: 09/18/2001 TIME: 11:35:00

Input Set : A:\sequence-text.txt

Output Set: N:\CRF3\09182001\I941179.raw

L:9 M:270 C: Current Application Number differs. Replaced Application Number

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:212 M:258 W: Mandatory Feature missing, <220> FEATURE: L:540 M:258 W: Mandatory Feature missing, <220> FEATURE:

L:869 M:258 W: Mandatory Feature missing, <220> FEATURE:

L:1366 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:75 SEQ:17

M:254 Repeated in SeqNo=17